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Minimum DB seq length: 0
Maximum DB seq length: 200000000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Database
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Perfect score:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Run on:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OM nucleic - nucleic search, using sw model
Published_Applications_NA:*

1: /cgn2_6/ptodata/2/pubpna/US07_PUBCOMB.seq:*

2: /cgn2_6/ptodata/2/pubpna/PCT_NEW_PUB.seq:*

3: /cgn2_6/ptodata/2/pubpna/US06_NEW_PUB.seq:*

4: /cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*

5: /cgn2_6/ptodata/2/pubpna/US07_NEW_PUB.seq:*

6: /cgn2_6/ptodata/2/pubpna/PCTUS_PUBCOMB.seq:*

7: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

9: /cgn2_6/ptodata/2/pubpna/US08_PUBCOMB.seq:*

9: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

10: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

11: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

12: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

13: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*

14: /cgn2_6/ptodata/2/pubpna/US09_NEW_PUB.seq:*
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Gapop 10.0 , Gapext 1.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  March 15, 2003, 15:05:45; Search time 16.1179 Seconds (without alignments) 10973.529 Million cell updates/sec
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AATTGTGCCCTTGAGGAGCT.....CCGGCAGCTGTGGCGCAGGA 252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GenCore version 5.1.4 p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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35.8	7 38.8	63			3 252					252	7 252	252	252	252	252	252	252	Score
14.2	15.4	25.0	69.0	98.8	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	100.0	Query Match
13500	5173	437	174	355	1996	1917	1917	1915	1915	1915	1915	626	626	626	563	563	563	Length
9 10	10	10	10	10	10	10	9	10	10	10	10	10	10	9	10	10	9	DB
US-09-923-876-4804 US-09-954-531-145	US-09-880-107-3356	US-09-998-598-2216	US-09-998-598-1740	US-09-867-701-4818	US-09-925-301-207	US-09-922-217-1105	US-10-025-380-1105	US-09-967-768A-192	US-09-880-107-3420	US-09-964-824A-563	US-09-964-824A-101	US-09-833-263-853	US-09-922-217-853	US-10-025-380-853	US-09-833-263-944	US-09-922-217-944	US-10-025-380-944	ID
Sequence 4804, A	Sequence 3356, P	Sequence 2216, A	Sequence 1740, A	Sequence 4818, A	Sequence 207, Ap	Sequence 1105, A	Sequence 1105, Ap	Sequence 192, App	Sequence 3420, A	Sequence 563, App	Sequence 101, App	Sequence 853, App	Sequence 853, App	Sequence 853, App	Sequence 944, App	Sequence 944, App	Sequence 944, App	Description

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32.4	2	32.6	32.6	32.6	32.6	32.6	32.6	32.6	32.6	32.6	32.8	33	ω ω	33.4	33.6	33.6	33.8	<b>ω</b>	<b>ω</b>	35.2	35.2	35.2	35.2	35.2	35.2
12.9	י פ	12.9	12.9	12.9	12.9	12.9	12.9	12.9	12.9	12.9	13.0	13.1	13.1	13.3		13.3	13.4	13.9	13.9		14.0	14.0	14.0	14.0	14.0
4055	35.40	2741	2740	2740	2740	2740	2740	1290	501	392	1680	10797	1041	4970	1214	687	25603	19616	2161	13500	13500	13500	13500	13500	13500
10	5	10	10	10	10	.10	10	10	10	9	10	10	10	10	10	10	9	10	10	10	10	10	10	10	œ
US-09-935-291A-1	IIC-09-935-2912-3	US-09-925-300-134	US-09-960-253-179	US-09-960-253-174	US-09-954-456-1158	US-09-954-456-710	US-09-954-456-77	US-09-815-242-7834	US-09-884-441-149	US-09-796-692-9300	US-09-815-242-7988	US-09-764-847-1577	US-09-815-242-7885	US-09-816-095-1	US-09-780-717-28	US-09-780-717-30	US-09-819-607-3	US-09-764-877-3220	US-09-810-808-4	US-09-880-107-2265	US-09-954-456-1801	US-09-954-456-1132	US-09-954-456-55	US-09-962-436-269	US-U9~954-531-363
Sequence 1, Appli	Semience 3 Appli	134	Sequence 179, App	Sequence 174, App	Sequence 1158, Ap	Sequence 710, App	Sequence 77, Appl	Sequence 7834, Ap	Sequence 149, App	Sequence 9300, Ap	Sequence 7988, Ap	Sequence 1577, Ap	Sequence 7885, Ap	Sequence 1, Appli	Sequence 28, Appl	Sequence 30, Appl	Sequence 3, Appli	Sequence 3220, Ap	Sequence 4, Appli	Sequence 2265, Ap	Sequence 1801, Ap	Sequence 1132, Ap	Sequence 55, Appl		Sequence 363, App

## ALIGNMENTS

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                                                                                                                                        ; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-025-380-944
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                                                                                                                                                                                           CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEG ID NOS: 1129
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 944
LENGTH: 563
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Xu, Jiangchun
                                                               Query Match
Best Local Similarity
Matches 252; Conserv
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APPLICANT:
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                                                      APPLICANT: Carter, Darrick TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE FILE REFERENCE: 210121.471C14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT:
472 AATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGGGACCAACTCCATGCC 413
                    1 AATTGTGCCCCTTGAGGAGCTGCGTCTGGGTCTTTGGGGCCTCTGGGGGGACCAACTCCATGCC 60
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Skeiky, Yasir A. W.
Fanger, Gary R.
Vedvick Thomas S.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Jiang, Yuqiu
Smith, Carole L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Stolk, John A. Wang, Tongtong
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Meagher, Madeleine Joy
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Benson, Darin R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Lodes, Michael J
                                                                   100.0%; Score 252; DB 9; ilarity 100.0%; Pred. No. 8.2e-59; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gordon E.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Heather
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                                                                                                     Length, 563;
                                                                     Indels
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                                                                   Gaps
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CURRENT APPLICANT: KU, Jiangchun
APPLICANT: Lodes, Michael J.
APPLICANT: Secrist, Heather
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Benson, Darin R.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Wang, Tongtong
APPLICANT: Wing, Gordon E.
APPLICANT: Wing, Gordon E.
APPLICANT: King, Gordon E.
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 944
LENGTH: 563
TYPE: DNA
ORGANISM: Homo sapiens
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US-09-922-217-944/c
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Best Local Similarity
Matches 252; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 AATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTGGGCCTCTGGGGGACCAACTCCATGCC 60
                                                                                                                                                                                                                                                                                   GAGAAGGATGGCCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGC
                                                                                                                                                                                                                                                                                                                           GAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCCAGGGCCCTTTGACCAGGGCAGC 180
                                                                                                                                                                                                                                                                                                                                                                                                                                  CAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG
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                                                                                                                                                                                AATTGTGCCCTTGAGGAGCTGCGTCTGGTCTTTTGGGCCCTCTGGGGGACCAACTCCATGCC 413
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                                                                  TGTGGCGCAGGA 252
                                                                                                                                           INFORMATION:
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   TGTGGCGCAGGA
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Pred. No. 8.2e-59;
); Mismatches 0;
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RESULT 3

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APPLICANT: Wang, Aijun
APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, John A.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Meagher, Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHER.
TITLE OF INVENTION: DIAGNOSIS OF COLON CANC.
FILE REFERENCE: 210121.471C12
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT FILING DATE: 2001-04-10
NUMBER OF SEQ ID NOS: 1093
SOFTWARE: FASTSEQ for Windows Version 3.0
SEQ ID NO 944
LENGTH: 563
TYPE: DNA
ORGANISM: Home sapien
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; Sequence 853, Application US/10025380
; Publication No. US20020182191A1
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                                                                                                                                                       APPLICANT: Xu, Jiangchun
APPLICANT: Lodes, Michael
APPLICANT: Secrist, Heath
APPLICANT: Secrist, Heath
APPLICANT: Benson, Darin
APPLICANT: Benson, Darin
APPLICANT: Meagher, Madel.
APPLICANT: Wang, Tongtong
APPLICANT: Wang, Yugiu
APPLICANT: Smith, Carole
APPLICANT: Smith, Carole
APPLICANT: King, Gordon E
APPLICANT: Wang, Aijun
APPLICANT: Clapper, Jonat.
APPLICANT: Clapper, Jonat.
APPLICANT: Shelky, Yasir
APPLICANT: Fangger, Gary R
APPLICANT: Vedvick Thomas
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Fanger, Gary R.
APPLICANT: Vedvick Thomas S.
APPLICANT: Carter, Darrick
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C14
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
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Stolk, John A.
Stolk, John A.
Wang, Tongtong
Jiang, Yugiu
Smith, Carole L.
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Wang, Aijun
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Benson, Darin R.
Meagher, Madeleine
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-025-380-853
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TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
CURRENT FILING DATE: 2001-08-03
NUMBER OF SEQ ID NOS: 1124
SOPTWARE: FRAESEQ for Windows Version 4.0
SEQ ID NO 853
LENGTH: 626
TYPE: DNA. 1670 626167
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APPLICANT: Lode
APPLICANT: Sec:
APPLICANT: Benn
APPLICANT: Mean
APPLICANT: Sto
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Best Local Similarity
Matches 252; Conserv
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SEQ ID NO 853
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Best Local Similarity
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                 CAGCTGCGAGACCTCACTTCCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG
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o. US20020076414A1
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Smith, Carole Lynn
King, Gordon E.
Wang, Aijun
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Benson, Darin R.
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Lodes, Michael J.
                                                                                                                    100.0%; Score 252; DB 10; ilarity 100.0%; Pred. No. 8.4e-59; Conservative 0; Mismatches 0;
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; Sequence 101, Application US/09964824A
; Patent No. US20020102531A1
; GENERAL INFORMATION:
; APPLICANT: HORRIGAN, Stephen
; TITLE OF INVENTION: Cancer Gene Determination
; TITLE OF INVENTION: Sets
; TITLE OF INVENTION: Sets
; TITLE REFERENCE: 689290-73
; CURRENT APPLICATION NUMBER: US/09/964,824A
; CURRENT FILING DATE: 2001-09-27
; PRIOR APPLICATION NUMBER: US/60/236,033
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SCOTUMARE: FastSEQ for Windows Version 3.0
SEQ ID NO 853
LENGTH: 626
TYPE: DNA
ORGANISM: Homo sapien
US-09-833-263-853
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US-09-964-824A-101
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Best Local Similarity
Matches 252; Conserv
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APPLICANT: Clapper, Jonathan D.
APPLICANT: Stolk, John A.
APPLICANT: Stolk, John A.
APPLICANT: Meagher, Madeleine J.
TITLE OF INVENTION: COMPOUNDS FOR COLON CANCER AND METALE REFERENCE: 210121.471012
CURRENT APPLICATION NUMBER: US/09/833,263
CURRENT FILING DATE: 2001-04-10
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PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 583
SOFTWARE: Patentin version 3.0
SEQ ID NO 101
LENGTH: 1915
TYPE: DNA
ORGANISM: Homo sapiens
US-09-964-824A-101
                                                                                                                                                                                                                                                                                     APPLICANT: HONORISM. Stephen
APPLICANT: HOVENTION: Cancer Gene Determination and
TITLE OF INVENTION: Sets
FILE REFERENCE: 689290-73
CURRENT APPLICATION NUMBER: US/69/964,824A
CURRENT FILING DATE: 2001-09-27
PRIOR APPLICATION NUMBER: US/60/236,033
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,032
PRIOR APPLICATION NUMBER: US/60/236,032
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PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR APPLICATION NUMBER: US/60/236,028
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEG ID NOS: 583
SOFTWARE: PatentIn version 3.0
SEG ID NO 563
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US-09-964-824A-563
J Sequence 563, Application US/09964824A
Patent No. US20020102531A1
Patent NoRMATION:
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Best Local Similarity 100.0%;
Matches 252; Conservative 0
                                                                                                                                       Matches 252; Conservative
                                                                                                                                                            Query Match
Best Local
                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                 ENGTH: 1915
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61 CAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG
                                                                                                                                                          Local Similarity
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Pred. No. 1.1e-58;
0; Mismatches 0;
                                                                                                                                       Score 252; DB 10;
Pred. No. 1.1e-58;
D; Mismatches 0;
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GENERAL INFORMATION:

APPLICANT: Horne, Darci T.

APPLICANT: Vockley, Joseph G.

APPLICANT: Scherf, Uwe

APPLICANT: Scherf, Uwe

APPLICANT: Scherf, Uwe

ITITLE OF INVENTION: Gene Expression Profiles in Liver Cancer

FILE REFERENCE: 44921-5028-WO

CURRENT APPLICATION NUMBER: US/09/880,107

CURRENT APPLICATION NUMBER: US/09/880,107

CURRENT FILING DATE: 2001-06-14

PRIOR APPLICATION NUMBER: US/09/87,054

PRIOR FILING DATE: 2000-10-02

NUMBER OF SEQ ID NOS: 3950

NUMBER OF SEQ ID NOS: 3950
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US-09-880-107-3420
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SEQ ID NO 3420
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                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: Genbank Accession No. US20020142981A1 U73843
                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
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                                                                                               GAGAAGGATGGCATGGCCTTCCAGGAGGCCCCTAGACCCAGGGCCCCTTTGACCAGGGCAGC
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RESULT 10
US-09-967-76BA-192
US-09-967-76BA-192
; Sequence 192, Application US,
; Patent No. US20020150877A1
; GENERAL INFORMATION:
; APPLICANT: Augustus, Meena

US/09967768A

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APPLICANT: Clapper, Jonathan D.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Skeiky, Yasir A. W.
APPLICANT: Stanger, Garry R.
APPLICANT: Vedvick Thomas S.
APPLICANT: Cartear, Darrick
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C1.4
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT APPLICATION NUMBER: US/10/025,380
CURRENT FILING DATE: 2001-12-19
NUMBER OF SEQ ID NOS: 1129
SOFTMARE: FastSEQ for Windows Version 4.0
SEQ ID NO 1105
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; ORGANISM: Homo sapiens
US-09-967-768A-192
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FILE REFERENCE: 689290-72
CURRENT APPLICATION NUMBER: US/09/967,768A
CURRENT FILING DATE: 2001-09-28
PRIOR APPLICATION NUMBER: US/60/236,109
PRIOR FILING DATE: 2000-09-28
PRIOR APPLICATION NUMBER: US/60/236,034
PRIOR APPLICATION NUMBER: US/60/236,034
PRIOR APPLICATION NUMBER: US/60/236,111
PRIOR APPLICATION NUMBER: US/60/236,111
PRIOR FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 325
SOFTWARE: PatentIn version 3.0
SEQ ID NO 192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 1105, Application US/10025380 Publication No. US20020182191A1
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Best Local Similarity
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Smith, Carole L.
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Benson, Darin R.
Meagher, Madeleine Joy
Stolk, John A.
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Lodes, Michael J
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; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 1105
; LENGTH: 1917
; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-922-217-1105
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; TYPE: DNA
; ORGANISM: Homo sapiens
US-10-025-380-1105
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                                                                                                                                                                            Query Match
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Best Local Similarity
                                                                                                                                                                          Matches 252;
                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Jiang, Yuqiu
APPLICANT: Smith, Carole Lynn
APPLICANT: Smith, Carole Lynn
APPLICANT: King, Gordon E.
APPLICANT: Wang, Aijun
APPLICANT: Clapper, Jonathan D.
APPLICANT: Clapper, Jonathan D.
TITLE OF INVENTION: COMPOUNDS FOR IMMUNOTHERAPY AND DIAGNOSIS
TITLE OF INVENTION: OF COLON CANCER AND METHODS FOR THEIR USE
FILE REFERENCE: 210121.471C13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               -09-922-217-1105
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                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/922,217
CURRENT FILING DATE: 2001-08-03
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121 GAGAAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGC 180
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                                                                  CAGCTGCGAGACCTCACTTCCAGCTCTTCTGATGAGCTCAGTTGGATCATTGAGCTGCTG 120
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Stolk, John A.
Wang, Tongtong
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Benson, Darin R.
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Pred. No. 1.1e-58;
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Pred. No. 1.1e-58
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APPLICANT: ROSEN et al.
TITLE OF INVENTION: Nucleic Acids, Proteins and Ant
FILE REFERENCE: PA106
CURRENT APPLICATION NUMBER: US/09/925,301
CURRENT FILING DATE: 2001-08-10
PRIOR APPLICATION NUMBER: PCT/US00/05882
PRIOR FILING DATE: 2000-03-08
PRIOR FILING DATE: 2000-03-08
PRIOR APPLICATION NUMBER: 60/124,270
PRIOR FILING DATE: 1999-03-12
NUMBER OF SEQ ID NOS: 1694
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 207
LENGTH: 1996
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US-09-925-301-207
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                                                                                                                                                                            Sequence 4818, Application US/09867701 Patent No. US20020132237A1
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Best Local
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ORGANISM: Homo sapiens
-09-925-301-207
APPLICANT: Aglate, Paul A.
APPLICANT: Jones, Robert
APPLICANT: Harlocker, Susan L.
APPLICANT: HARLOCKER, SUSAN L.
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY
TITLE OF INVENTION: AND DIAGNOSIS OF OVARIAN CANCER
FILE REFERENCE: 210121.497
CURRENT APPLICATION NUMBER: US/09/867,701
CURRENT FILING DATE: 2001-05-29
NUMBER OF SEQ ID NOS: 10912
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                                                                                                                                                                                                                                                                                                                                                       tch 100.0%; Score 252; DB 10; al Similarity 100.0%; Pred. No. 1.1e-58; 252; Conservative 0; Mismatches 0;
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APPLICANT: Stolk, John A.
APPLICANT: Xu, Jiangchun
APPLICANT: Chenault, Ruth A.
APPLICANT: Meagher, Madelein Joy
APPLICANT: Meagher, Madelein Joy
TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR THE THERAPY AND
TITLE OF INVENTION: DIAGNOSIS OF COLON CANCER
FILE REFERENCE: 210121.561
CURRENT APPLICATION NUMBER: US/09/98,598
CURRENT FILING DATE: 2001-11-16
NUMBER OF SEQ ID NOS: 2606
SOFTWARE: COTIXA Invention Disclosure Database
SEQ ID NO 1740
LENGTH: 174
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Best Local S
Matches 249
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SEQ ID NO 4818
LENGTH: 355
                                                                                                                                                                                                                      Query Match 69.0%; Score 174; DB 10; Best Local Similarity 100.0%; Pred. No. 5.4e-38; Matches 174; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1740, Application US/09998598 Patent No. US20020150922A1
                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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Local Similarity 100.0%; Pred. No. 4.6e-58;
hes 249; Conservative 0; Mismatches 0;
                                                                         114 AAGGATGGCATGGCCTTCCAGGAGGCCCTAGACCCAGGGCCCTTTGACCAGGGCAGCCCC
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355
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